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<110> WANG, XIAODONG
LIU, XUESONG

<120> DNA FRAGMENTATION FACTOR INVOLVED IN APOPTOSIS

<130> UTSD:546USD1

<140> UNKNOWN

<141> 2000-12-22

<150> 09/061,702

<151> 1998-04-16

<160> 21

<170> PatentIn Ver. 2.1

<210> 1

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<213> Homo sapiens

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 Tyr Glu Asp Gly Thr Glu Leu Thr Glu Asp Tyr Phe Pro Ser Val Pro
 50 55 60
 Asp Asn Ala Glu Leu Val Leu Leu Thr Leu Gly Gln Ala Trp Gln Gly
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002233T-133200

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Pro	Gln	Arg	Gln	Arg	Leu	Leu	Ala	Asp	Leu	Leu	His	Asn	Val	Ser	Gln	115	120	125	
Asn	Ile	Ala	Ala	Glu	Thr	Arg	Ala	Glu	Asp	Pro	Pro	Trp	Phe	Glu	Gly	130	135	140	
Leu	Glu	Ser	Arg	Phe	Gln	Ser	Lys	Ser	Gly	Tyr	Leu	Arg	Tyr	Ser	Cys	145	150	155	160
Glu	Ser	Arg	Ile	Arg	Ser	Tyr	Leu	Arg	Glu	Val	Ser	Ser	Tyr	Pro	Ser	165	170	175	
Thr	Val	Gly	Ala	Glu	Ala	Gln	Glu	Glu	Phe	Leu	Arg	Val	Leu	Gly	Ser	180	185	190	
Met	Cys	Gln	Arg	Leu	Arg	Ser	Met	Gln	Tyr	Asn	Gly	Ser	Tyr	Phe	Asp	195	200	205	
Arg	Gly	Ala	Lys	Gly	Gly	Ser	Arg	Leu	Cys	Thr	Pro	Glu	Gly	Trp	Phe	210	215	220	
Ser	Cys	Gln	Gly	Pro	Phe	Asp	Met	Asp	Ser	Cys	Leu	Ser	Arg	His	Ser	225	230	235	240
Ile	Asn	Pro	Tyr	Ser	Asn	Arg	Glu	Ser	Arg	Ile	Leu	Phe	Ser	Thr	Trp	245	250	255	
Asn	Leu	Asp	His	Ile	Ile	Glu	Lys	Lys	Arg	Thr	Ile	Ile	Pro	Thr	Leu	260	265	270	
Val	Glu	Ala	Ile	Lys	Glu	Gln	Asp	Gly	Arg	Glu	Val	Asp	Trp	Glu	Tyr	275	280	285	
Phe	Tyr	Gly	Leu	Leu	Phe	Thr	Ser	Glu	Asn	Leu	Lys	Leu	Val	His	Ile	290	295	300	
Val	Cys	His	Lys	Lys	Thr	Thr	His	Lys	Leu	Asn	Cys	Asp	Pro	Ser	Arg	305	310	315	320
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<211> 1689

<212> DNA

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Met Glu

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gtg acc ggg gac gcc ggg gta cca gaa tct ggc gag atc cgg act cta 166
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aag ccg tgt ctg ctg cgc cgc aac tac agc cgc gaa cag cac ggc gtg 214
Lys Pro Cys Leu Leu Arg Arg Asn Tyr Ser Arg Glu Gln His Gly Val
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gcc gcc tcc tgc ctc gaa gac ctg agg agc aag gcc tgt gac att ctg 262
Ala Ala Ser Cys Leu Glu Asp Leu Arg Ser Lys Ala Cys Asp Ile Leu
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gcc att gat aag tcc ctg aca cca gtc acc ctt gtc ctg gca gag gat 310
Ala Ile Asp Lys Ser Leu Thr Pro Val Thr Leu Val Leu Ala Glu Asp
55 60 65

ggc acc ata gtg gat gat gac gat tac ttt ctg tgt cta cct tcc aat 358
Gly Thr Ile Val Asp Asp Asp Asp Tyr Phe Leu Cys Leu Pro Ser Asn
70 75 80

act aag ttt gtg gca ttg gct agt aat gag aaa tgg gca tac aac aat 406
Thr Lys Phe Val Ala Leu Ala Ser Asn Glu Lys Trp Ala Tyr Asn Asn
85 90 95

tca gat gga ggt aca gct tgg att tcc caa gag tcc ttt gat gta gat 454
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100 105 110

002221.15484260

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Leu	Lys	Glu	Asp	Leu	Ser	Ser	Ile	Ile	Leu	Leu	Ser	Glu	Glu	Asp	Leu	
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cag	atg	ctt	gtt	gac	gct	ccc	tgc	tca	gac	ctg	gct	cag	gaa	cta	cgt	598
Gln	Met	Leu	Val	Asp	Ala	Pro	Cys	Ser	Asp	Leu	Ala	Gln	Glu	Leu	Arg	
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Gln	Ser	Cys	Ala	Thr	Val	Gln	Arg	Leu	Gln	His	Thr	Leu	Gln	Gln	Val	
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Tyr	Leu	Gln	Ala	Leu	Glu	Lys	Glu	Gly	Ser	Leu	Leu	Ser	Lys	Gln	Glu	
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Glu	Ser	Lys	Ala	Ala	Phe	Gly	Glu	Glu	Val	Asp	Ala	Val	Asp	Thr	Gly	
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act	gca	ctg	agg	gag	aag	cag	gct	cca	gag	ctg	agc	tta	tct	agt	cag	886
Thr	Ala	Leu	Arg	Glu	Lys	Gln	Ala	Pro	Glu	Leu	Ser	Leu	Ser	Ser	Gln	
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Asp	Leu	Glu	Leu	Val	Thr	Lys	Glu	Asp	Pro	Lys	Ala	Leu	Ala	Val	Ala	
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Leu	Asn	Trp	Asp	Ile	Lys	Lys	Thr	Glu	Thr	Val	Gln	Glu	Ala	Cys	Glu	
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 Pro Lys Arg Ala Arg Gln Asp Pro Thr
 325 330

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<213> Homo sapiens

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 35 40 45

Ile Leu Ala Ile Asp Lys Ser Leu Thr Pro Val Thr Leu Val Leu Ala
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Glu Asp Gly Thr Ile Val Asp Asp Asp Tyr Phe Leu Cys Leu Pro

002222T#TSH260

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Val Asp Glu Thr Asp Ser Gly Ala Gly Leu Lys Trp Lys Asn Val Ala						
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Arg Glu Leu Lys Glu Asp Leu Ser Ser Ile Ile Leu Leu Ser Glu Glu						
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Asp Leu Gln Met Leu Val Asp Ala Pro Cys Ser Asp Leu Ala Gln Glu						
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Ile Leu Thr Ala Leu Arg Glu Lys Gln Ala Pro Glu Leu Ser Leu Ser						
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00222T "TSH260

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<223> D = G, C or A

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00343454-122400

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<400> 21
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1 5

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